

SEQUENCE LISTING

<110> Ben-Sasson, Shmuel A.

<120> Short Peptides Which Selectively
Modulate the Activity of Protein Kinases

<130> 1242.1029-000 (CMCC-679)

<140> US 09/161,094

<141> 1998-09-25

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Gly Arg Phe

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Gly Arg Phe

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 Gly Val Phe

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Asp Glu Asp Asn Pro Gly Phe
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Leu Thr Ile Met Asn Gly Gly Asp Leu Lys Phe His Ile Tyr Asn Met
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Gly Asn Pro Gly Phe
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Gly Gln Ala Gly Phe
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 Gly Gly Tyr

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 Lys Ala Leu

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 Lys Ala Val

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 Val Lys Pro Phe
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 His Thr Leu

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Lys Gln Leu

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Gly Arg Met

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 Gly Arg Met

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Leu Thr Gly Ile
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Pro Pro Gly Leu
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Thr Gly Lys Tyr Leu
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 Asp Gly Lys Tyr Leu
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 Glu Gly Gln Asp Leu
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 Glu Gly Gly Lys Val
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12/55

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1 5 10 15
Glu Gly Ser Lys Gln
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Ser Gly Ile Lys Leu
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Gly Arg Ser Val Leu
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1 5 10 15
Gly Arg Ala Leu Val
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Lys Tyr Ser Leu
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Asn Arg Gln Glu Val
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Arg Val Leu Glu Thr Asp Pro Ala Phe Ala Arg Glu His Gly Thr Ala
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Ser Thr Leu
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<400> 50
Ile Glu Tyr Ala Pro His Gly Asn Leu Leu Asp Phe Leu Arg Lys Ser
1 5 10 15

Arg Val Leu Glu Thr Asp Pro Ala Phe Ala Ile Ala Asn Ser Thr Ala
 20 25 30
 Ser Thr Leu
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<400> 51
 Val Glu Tyr Ala Ser Lys Gly Asn Leu Arg Glu Tyr Leu Gln Ala Arg
 1 5 10 15
 Arg Pro Pro Gly Leu Glu Tyr Cys Tyr Asn Pro Ser His Asn Pro Glu
 20 25 30
 Glu Gln Leu
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<400> 52
 Val Glu Tyr Ala Ser Lys Gly Asn Leu Arg Glu Tyr Leu Arg Ala Arg
 1 5 10 15
 Arg Pro Pro Gly Met Glu Tyr Ser Tyr Asp Ile Asn Arg Val Pro Glu
 20 25 30
 Glu Gln Met
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 Val Glu Tyr Ala Ala Lys Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg
 1 5 10 15
 Arg Pro Pro Gly Leu Asp Tyr Ser Phe Asp Thr Cys Lys Pro Pro Glu
 20 25 30
 Glu Gln Leu
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 Val Glu Cys Ala Ala Lys Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg
 1 5 10 15
 Arg Pro Pro Gly Pro Asp Leu Ser Pro Asp Gly Pro Arg Ser Ser Glu
 20 25 30
 Gly Pro Leu
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 1 5 10 15
 Arg Asp Ser Phe Leu Ser His His Pro Glu Lys Pro Lys Lys Glu Leu
 20 25 30
 Ile Phe Gly Leu Asn Pro Ala
 35 40

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 Thr Glu Tyr Cys Arg Tyr Gly Asp Leu Val Asp Tyr Leu His Arg Asn
 1 5 10 15
 Lys His Thr Phe Leu Gln His His Ser Asp Lys Arg Arg Pro Pro Ser
 20 25 30
 Ala Glu Leu Tyr Ser Asn Ala Leu
 35 40

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 Val Glu Tyr Cys Lys Tyr Gly Asn Leu Ser Asn Tyr Leu Lys Ser Lys
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 Arg Asp Leu Phe Phe Leu Asn Lys Asp Ala Ala Leu His Met Glu Pro
 20 25 30
 Lys Lys Glu Lys Met Glu Pro Gly
 35 40

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Val Glu Phe Cys Lys Tyr Gly Asn Leu Ser Asn Phe Leu Arg Ala Lys
 1 5 10 15
 Arg Asp Ala Phe Ser Pro Cys Ala Glu Lys Ser Pro Glu Gln Arg Gly
 20 25 30
 Arg Phe Arg Ala Met Val Glu Leu
 35 40

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Val Glu :

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Leu Pro Tyr Met
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Thr His Asn Pro
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Leu Pro Tyr Met Arg His Gly Asp Leu Arg His Phe Ile Arg Ala Gln
1 5 10 15
Glu Arg Ser Pro
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1 5 10 15
Gln Arg Asn Pro
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Thr Gln Leu Met Pro Phe Gly Cys Leu Leu Asp Tyr Val Arg Glu His
1 5 10 15
Lys Asp Asn Ile
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Arg Gly Arg Leu

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Arg Gly Ala Leu
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1 5 10 15
Lys Asp Asn Ile
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Val Glu Tyr Ala Lys Tyr Gly Ser Leu Arg Gly Phe Leu Arg Glu Ser
1 5 10 15
Arg Lys Val Gly Pro Gly Tyr Leu Gly Ser Gly Ser Arg Asn Ser
20 25 30
Ser Ser Leu Asp His Pro Asp Glu Arg Ala Leu
35 40

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 Gly Pro Asp Ala Lys Leu Leu Ala Gly Gly Glu Asp Val Ala Pro Gly
 20 25 30
 Pro Leu.

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 1 5 10 15
 Gly Pro Asp Ala Val Leu Met Ala Glu Gly Asn Pro Pro Thr Glu Leu
 20 25 30

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 Gly Pro Asp Ala Met Ile Leu Val Asp Gly Gln Pro Arg Gln Ala Lys
 20 25 30
 Gly Glu Leu
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 Met Glu Met Ala Glu Leu Gly Pro Leu Asn Lys Tyr Leu Gln Gln Asn
 1 5 10 15
 Arg His Val

<210> 72

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<400> 72
Met Glu Met Ala Gly Gly Pro Leu His Lys Phe Leu Val Gly Lys
1 5 10 15
Arg Glu Glu Ile
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<210> 73
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Met Glu Phe Leu Pro Ser Gly Ser Leu Lys Glu Tyr Leu Pro Lys Asn
1 5 10 15

Lys Asn Lys Ile
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Met Glu Tyr Leu Pro Tyr Gly Ser Leu Arg Asp Tyr Leu Gln Lys His
1 5 10 15
Lys Glu Arg Ile
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1 5 10 15
Arg Ala Arg Leu
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1 5 10 15
Ser Ile

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Leu	Glu	Tyr	Ala	Pro	Leu	Gly	Thr	Val	Tyr	Arg	Glu	Leu	Gln	Lys	Leu
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Ser Lys Phe

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Leu	Glu	Tyr	Cys	Ser	Gly	Gly	Glu	Leu	Phe	Asp	Arg	Ile	Glu	Pro	Asp
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Ile Gly Met

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Met	Glu	Tyr	Cys	Ser	Gly	Gly	Asp	Leu	Arg	Lys	Leu	Leu	Asn	Lys	Pro
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Glu Asn Cys Cys Gly Leu

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Met	Glu	Tyr	Cys	Gln	Gly	Gly	Asp	Leu	Arg	Lys	Tyr	Leu	Asn	Gln	Phe
1					5			10						15	

Glu Asn Cys Cys Gly Leu

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<211> 19

23/55

<212> PRT
<213> unknown

<220>
<223> DAPK

<400> 81
Leu Glu Leu Val Ala Gly Gly Glu Leu Phe Asp Phe Leu Ala Glu Lys
1 5 10 15
Glu Ser Leu

<210> 82
<211> 31
<212> PRT
<213> unknown

<220>
<223> IRK

<400> 82
Met Glu Leu Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu
1 5 10 15
Arg Pro Glu Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu
20 25 30

<210> 83
<211> 18
<212> PRT
<213> unknown

<220>
<223> TGFbRII

<400> 83
Thr Ala Phe His Ala Lys Gly Asn Leu Gln Glu Tyr Leu Thr Arg His
1 5 10 15
Val Ile

<210> 84
<211> 18
<212> PRT
<213> unknown

<220>
<223> ACTRIIA

<400> 84
Thr Ala Phe His Glu Lys Gly Ser Leu Ser Asp Phe Leu Lys Ala Asn
1 5 10 15
Val Val

<210> 85
<211> 18
<212> PRT
<213> unknown

<220>
<223> ACTRIIB

<400> 85
Thr Ala Phe His Asp Lys Gly Ser Leu Thr Asp Tyr Leu Lys Gly Asn
1 5 10 15
Ile Ile

<210> 86
<211> 18
<212> PRT
<213> unknown

<220>
<223> ALK1

<400> 86
Thr His Tyr His Glu His Gly Ser Leu Tyr Asp Phe Leu Gln Arg Gln
1 5 10 15
Thr Leu

<210> 87
<211> 18
<212> PRT
<213> unknown

<220>
<223> ALK2

<400> 87
Thr His Tyr His Glu Met Gly Ser Leu Tyr Asp Tyr Leu Gln Leu Thr
1 5 10 15
Thr Leu

<210> 88
<211> 18
<212> PRT
<213> unknown

<220>
<223> ALK3

<400> 88
Thr Asp Tyr His Glu Asn Gly Ser Leu Tyr Asp Phe Leu Lys Cys Ala

1 5 10 15
Thr Leu

<210> 89
<211> 18
<212> PRT
<213> unknown

<220>
<223> ALK4

<400> 89
Ser Asp Tyr His Glu His Gly Ser Leu Phe Asp Tyr Leu Asn Arg Tyr
1 5 10 15

Thr Val

CD

<210> 90
<211> 18
<212> PRT
<213> unknown

<220>
<223> alk6

<400> 90
Thr Asp Tyr His Glu Asn Gly Ser Leu Tyr Asp Tyr Leu Lys Ser Thr
1 5 10 15
Thr Leu

<210> 91
<211> 18
<212> PRT
<213> unknown

<220>
<223> DDR1

<400> 91
Thr Asp Tyr Met Glu Asn Gly Asp Leu Asn Gln Phe Leu Ser Ala His
1 5 10 15
Gln Leu

<210> 92
<211> 18
<212> PRT
<213> unknown

<220>
<223> DDR2

<400> 92
Thr Glu Tyr Met Glu Asn Gly Asp Leu Asn Gln Phe Leu Ser Arg His
1 5 10 15
Glu Pro

<210> 93
<211> 21
<212> PRT
<213> unknown

<220>
<223> ILK
<400> 93

Thr His Trp Met Pro Tyr Gly Ser Leu Tyr Asn Val Leu His Glu Gly
1 5 10 15
Thr Asn Phe Val Val
20

<210> 94
<211> 16
<212> PRT
<213> unknown

<220>
<223> JNK

<400> 94
Met Glu Leu Met Asp Ala Asn Leu Cys Gln Val Ile Gln Met Glu Leu
1 5 10 15

<210> 95
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)
<223>

<221> AMIDATION
<222> (0)...(20)

<223> Akt1/Raca

<400> 95
Gly Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser Arg
1 5 10 15
Glu Arg Val Phe
20

<210> 96
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(19)

<223> Alk1

<400> 96
Gly Thr His Tyr His Glu His Gly Ser Leu Tyr Asp Phe Leu Gln Arg
1 5 10 15

Gln Thr Leu

<210> 97
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

~~B~~ <223> Braf

~~B~~ <400> 97
Lys Lys Lys Lys Lys Lys Gly Gly Ser Ser Leu Tyr His His Leu His
1 5 10 15
Ile Ile Glu Thr Lys Phe
20

~~B~~ <210> 98
<211> 21
<212> PRT
<213> Artificial Sequence

~~C~~ <220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(21)

<223> Braf

<400> 98
Gly Thr Gln Trp Ser Glu Gly Ser Ser Leu Tyr His His Leu His Ile
1 5 10 15
Ile Glu Thr Lys Phe
20

<210> 99
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> c-Ab1

<400> 99
 Gly Thr Glu Phe Met Thr Tyr Gly Asn Leu Leu Asp Tyr Leu Arg Glu
 1 5 10 15
 Cys Asn Arg Gln Glu Val
 20

<210> 100
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
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 <222> (1)...(0)

H<221> AMIDATION
 S<222> (0)...(21)
 D<223>

S

S

S<223> c-Met

SP

SP<400> 100

Gly Leu Pro Tyr Met Lys His Gly Asp Leu Arg Asn Phe Ile Arg Asn
 1 5 10 15
 Glu Thr His Asn Pro
 20

SP

SP

SP<210> 101

SP<211> 21

SP<212> PRT

<213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)

<223> c-Raf

<400> 101

Gly Thr Gln Trp Ser Glu Gly Ser Ser Leu Tyr Lys His Leu His Val
 1 5 10 15
 Gln Glu Thr Lys Phe
 20

<210> 102
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)

<223> benzyl ester at position 11

<221> AMIDATION

<222> (0)...(14)

<223> c-Raf

<400> 102

Ser	Ser	Leu	Tyr	Lys	His	Leu	His	Val	Gln	Glu	Thr	Lys	Phe
1						5			10				

<210> 103

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(21)

<223> c-Sea

<400> 103

Gly	Leu	Pro	Tyr	Met	Arg	His	Gly	Asp	Leu	Arg	His	Phe	Ile	Arg	Ala
1				5					10				15		
Gln	Glu	Arg	Ser	Pro											
				20											

<210> 104

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223> c-Src

<400> 104

Gly	Thr	Glu	Tyr	Met	Ser	Lys	Gly	Ser	Leu	Leu	Asp	Phe	Leu	Lys	Gly
1					5				10				15		
Glu	Thr	Gly	Lys	Tyr	Leu										
				20											

<210> 105

<211> 14

<212> PRT

<213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 5
 benzyl ester at position 9

<221> AMIDATION
 <222> (0)...(14)

<223> c-Src

<400> 105
 Gly Ser Leu Leu Asp Leu Lys Gly Glu Thr Gly Lys Phe Leu
 1 5 10

<210> 106
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)
 <223>

<223> CDK2

<400> 106
 Gly Phe Glu Phe Leu His Gln Asp Leu Lys Lys Phe Met Asp Ala Ser
 1 5 10 15
 Ala Leu Thr Gly Ile
 20

<210> 107
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 1
 benzyl ester at position 7

<221> AMIDATION
 <222> (0)...(14)
 <223>

<223> CDK2

<400> 107

Asp Leu Lys Lys Phe Met Asp Ala Ser Ala Leu Thr Gly Met
1 5 10

<210> 108
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1) ... (0)
<223> benzyl ester at position 1
benzyl ester at position 7

<221> AMIDATION
<222> (0) ... (14)

<223> CDK4

<400> 108

Asp Leu Arg Thr Tyr Leu Asp Lys Ala Pro Pro Pro Gly Leu
1 5 10

<210> 109
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1) ... (0)

<221> AMIDATION
<222> (0) ... (21)

<223> CDK4

<400> 109

Gly Phe Glu His Val Asp Gln Asp Leu Arg Thr Tyr. Leu Asp Lys Ala
1 5 10 15
Pro Pro Pro Gly Leu
20

<210> 110
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1) ... (0)

<221> AMIDATION
<222> (0) ... (21)

<223> CDK6

<400> 110
Gly Phe Glu His Val Asp Gln Asp Leu Thr Thr Tyr Leu Asp Lys Val
1 5 10 15
Pro Glu Pro Gly Val
20

<210> 111
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(19)

<223> Chk1

<400> 111
Gly Glu Tyr Ser Ser Gly Gly Glu Leu Phe Asp Arg Ile Glu Pro Asp
1 5 10 15
Ile Gly Met

<210> 112
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(19)
<223>

<223> Chk1

<400> 112
Gly Glu Tyr Ala Ser Gly Gly Glu Leu Phe Asp Arg Ile Glu Pro Asp
1 5 10 15
Ile Gly Met

<210> 113
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
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<222> (1)...(0)

<221> AMIDATION
<222> (0)...(19)
<223>

<223> CK IIa

<400> 113
Lys Lys Lys Lys Lys Gly Gly Asn Asn Thr Asp Phe Lys Gln Leu Tyr
1 5 10 15
Gln Thr Leu

<210> 114
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(17)

<223> CK IIa

<400> 114
Gly Phe Glu His Val Asn Asn Thr Asp Phe Lys Gln Leu Tyr Gln Thr
1 5 10 15
Leu

<210> 115
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)
<223>

<223> Csk

<400> 115
Gly Thr Glu Tyr Met Ala Lys Gly Ser Leu Val Asp Tyr Leu Arg Ser
1 5 10 15
Arg Gly Arg Ser Val Leu
20

<210> 116

<211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 5

<221> AMIDATION
 <222> (0)...(14)

<223> Csk

<400> 116
 Gly Ser Leu Val Asp Leu Arg Ser Arg Gly Arg Ser Val Leu
 1 5 10

<210> 117
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)

<223> Fak

<400> 117
 Gly Met Glu Leu Ser Thr Leu Gly Glu Leu Arg Ser Phe Leu Gln Val
 1 5 10 15
 Arg Lys Tyr Ser Leu
 20

<210> 118
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(17)

<223> FGFR-3

<400> 118
 Gly Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Arg Pro Pro Gly Leu
 1 5 10 15
 Glu

<210> 119
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1) ... (0)
<223> benzyl ester at position 5
benzyl ester at position 16

<221> AMIDATION
<222> (0) ... (16)

<223> FGFR-3

<400> 119
Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Pro Pro Gly Leu Glu
1 5 10 15

<210> 120
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1) ... (0)
<221> AMIDATION
<222> (0) ... (23)

<223> FGFR-3

<400> 120
Gly Val Glu Tyr Ala Ala Lys Gly Asn Leu Arg Glu Phe Leu Arg Ala
1 5 10 15
Arg Arg Pro Pro Gly Leu Glu
20

<210> 121
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> stearyl at position 1

<221> AMIDATION
<222> (0) ... (13)
<223> FGFR-3

<400> 121
Gly Ser Phe Asp Thr Ser Lys Pro Pro Glu Glu Gln Leu
1 5 10

<210> 122
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(23)

<223> Flk1

<400> 122
Gly Val Glu Phe Ser Lys Phe Gly Asn Leu Ser Asn Phe Leu Arg Ala
1 5 10 15
Lys Arg Asn Leu Phe Val Pro
20

<210> 123
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(17)
<223>

<223> Flk1

<400> 123
Gly Gly Asn Leu Ser Asn Phe Leu Arg Ala Lys Arg Asn Leu Phe Val
1 5 10 15
Pro

<210> 124
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(16)

<223> Flk1

<400> 124
Gly Asn Leu Ser Asn Phe Leu Arg Ala Lys Arg Asn Leu Phe Val Pro
1 5 10 15

<210> 125
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> stearyl at position 1

<221> AMIDATION
<222> (0)...(13)
<223> Flk1

<400> 125
Gly Arg Phe Arg Gln Gly Lys Asp Tyr Val Gly Glu Leu
1 5 10

<210> 126
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)
<221> AMIDATION
<222> (0)...(22)
<223> GSK3b

<400> 126
Lys Lys Lys Lys Lys Gly Gly Val Ala Arg His Tyr Ser Arg
1 5 10 15
Ala Lys Gln Thr Leu Pro
20

<210> 127
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)
<221> AMIDATION
<222> (0)...(13)
<223> GSK3b

<400> 127
Val Ala Arg His Tyr Ser Arg Ala Lys Gln Thr Leu Pro
1 5 10

<210> 128
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> GSK3b

<400> 128

Gly Asp Tyr Val Pro Glu Thr Val Tyr Arg Val Ala Arg His Tyr Ser
1 5 10 15
Arg Ala Lys Gln Thr Leu
20

<210> 129
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(12)

L <223> GSK3b

129
Arg Val Ala Arg His Tyr Ser Arg Ala Lys Gln Thr
1 5 10

130
<210> 130
<211> 22
<212> PRT
<213> Artificial Sequence

130
<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> Hck

130
Gly Thr Glu Phe Met Ala Lys Gly Ser Leu Leu Asp Phe Leu Lys Ser
1 5 10 15
Asp Glu Gly Ser Lys Gln
20

<210> 131
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
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<222> (1)...(0)

<221> AMIDATION

<222> (0) . . . (20)

<223> Tak1

<400> 131
 Gly Leu Glu Tyr Ala Pro Leu Gly Thr Val Tyr Arg Glu Leu Gln Lys
 1 5 10 15
 Leu Ser Lys Phe
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<210> 132

<211> 23

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1) . . . (0)

(2, 3, 4, 5, 6)

<221> AMIDATION

<222> (0)...(23)

6

<400> 132

Gly Met Glu Tyr Ser Ser Gly Gly Asp Leu Arg Lys Leu Leu Asn Lys
 1 5 10 15
 Pro Glu Asn Ser Ser Gly Leu
 20

<210> 133

<211> 23

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1) . . . (0)

<221> AMIDATION

<222> (0) . . . (23)

<223>

<223> IKK-2

<400> 133

Gly Met Glu Tyr Ser Gln Gly Gly Asp Leu Arg Lys Tyr Leu Asn Gln
 1 5 10 15
 Phe Glu Asn Ser Ser Gly Leu
 20

<210> 134
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
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<221> AMIDATION
<222> (0)...(22)

<223> ILK

~~134~~ <400> 134
Gly Thr His Trp Met Pro Tyr Gly Ser Leu Tyr Asn Val Leu His Glu
1 5 10 15
Gly Thr Asn Phe Val Val
20

<210> 135
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> stearyl at position 1

<221> AMIDATION
<222> (0)...(13)
<223> ILK

<400> 135
Gly Tyr Asn Val Leu His Glu Gly Thr Asn Phe Val Val
1 5 10

<210> 136
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(19)
<223>

<223> IRK

<400> 136
Gly Met Glu Leu Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser
1 5 10 15
Leu Arg Pro

<210> 137
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(12)

<223> IRK

<400> 137
Ala Gln Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu
1 5 10

<210> 138
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> IRK

<400> 138
Gly Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu Ala
1 5 10

<210> 139
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
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<222> (1)...(0)

<221> AMIDATION
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<223> IRK

<400> 139
Gly Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu
1 5 10

<210> 140
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
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<222> (1)...(0)

<221> AMIDATION
<222> (0)...(17)

<223> IRK

<400> 140
Gly Leu Arg Pro Glu Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr
1 5 10 15
Leu

<210> 141
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(21)

<223> Jak1

<400> 141
Gly Met Glu Phe Leu Pro Ser Gly Ser Leu Lys Glu Tyr Leu Pro Lys
1 5 10 15
Asn Lys Asn Lys Ile
20

<210> 142
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> Jak1

<400> 142
Gly Leu Lys Glu Tyr Leu Pro Lys Asn Lys Asn Lys Ile
1 5 10

<210> 143
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> Jak2

<400> 143
Gly Leu Arg Asp Tyr Leu Gln Lys His Lys Glu Arg Ile
1 5 10

<210> 144
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> stearyl at position 1

<221> AMIDATION
<222> (0)...(11)
<223> Jak2

<400> 144
Gly Leu Arg Asp Tyr Leu Gln Lys His Lys Glu
1 5 10

<210> 145
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(20)

<223> Jak3

<400> 145

Gly Met Glu Tyr Leu Pro Ser Gly Ser Leu Arg Asp Phe Leu Gln Arg
1 5 10 15
His Arg Ala Leu
20

<210> 146

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(21)

<223> Jak3

<400> 146

Gly Met Glu Tyr Leu Pro Ser Gly Ser Leu Arg Asp Phe Leu Gln Arg
1 5 10 15
His Arg Ala Arg Leu
20

<210> 147

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(13)

<223> Jak3

<400> 147

Gly Leu Arg Asp Phe Leu Gln Arg His Arg Ala Arg Leu
1 5 10

<210> 148

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<221> ACETYLATION

<222> (1)...(0)
<223> benzyl ester at position 5

<221> AMIDATION
<222> (0)...(14)

<223> Lck

<400> 148
Gly Ser Leu Val Asp Leu Lys Thr Pro Ser Gly Ile Lys Leu
1 5 10

<210> 149
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> Lck

<400> 149
Gly Thr Glu Tyr Met Glu Asn Gly Ser Leu Val Asp Phe Leu Lys Thr
1 5 10 15
Pro Ser Gly Ile Lys Leu
20

<210> 150
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> Lyn

<400> 150
Gly Thr Glu Tyr Met Ala Lys Gly Ser Leu Leu Asp Phe Leu Lys Ser
1 5 10 15
Asp Glu Gly Gly Lys Val
20

<210> 151
<211> 20
<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE
<222> (1) . . . (0)

<221> AMIDATION
<222> (0) . . . (20)

<223> MARK1

<400> 151

Gly Met Glu Tyr Ala Ser Gly Gly Glu Val Phe Asp Tyr Leu Val Ala
1 5 10 15
His Gly Arg Met
20

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15 <210> 152

212 PBT

<213> Artificial Sequence

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~~220~~ <221> ACETYLATION

~~222~~ <222> ACETYLATE
<222> (1)...(0)

<223> benzyl ester at position 2
benzyl ester at position 5

<221> AMIDATION

<222> (0) . . . (15)

<223> PDGFR-b

 U.S. GOVERNMENT PRINTING OFFICE 152

Gly Asp Leu Val Asp Tyr Leu His Arg Asn Lys His Thr Phe Leu
 1 5 10 15

<210> 153

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1) . . . (0)

<221> AMIDATION

<222> (0) . . . (22)

<223> PDGFR-b

<400> 153

Gly Thr Glu Tyr Ser Arg Tyr Gly Asp Leu Val Asp Tyr Leu His Arg
 1 5 10 15
 Asn Lys His Thr Phe Leu
 20

<210> 154
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
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<223> PKC_b

<400> 154
Gly Met Glu Tyr Val Asn Gly Gly Asp Leu Met Tyr His Ile Gln Gln
1 5 10 15
Val Gly Arg Phe
20

<210> 155
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)
<221> AMIDATION
<222> (0)...(20)

<223> PKC_b

<400> 155
Lys Lys Lys Lys Lys Gly Gly Asp Leu Met Tyr His Ile Gln Gln
1 5 10 15
Val Gly Arg Phe
20

<210> 156
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)
<223> benzyl ester at position 5

<221> AMIDATION
<222> (0)...(12)

<223> Plk

<400> 156
Arg Ser Leu Leu Glu Leu His Lys Arg Arg Lys Ala

1

5

10

<210> 157
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)
<223> benzyl ester at position 6

<221> AMIDATION
<222> (0)...(13)

<223> Plk

157
Gly Arg Ser Leu Leu Glu Leu His Lys Arg Arg Lys Ala
1 5 10

158
<210> 158
<211> 20
<212> PRT
<213> Artificial Sequence

158
<220>
<221> MYRISTATE
<222> (1)...(0)
AMIDATION
<221> AMIDATION
<222> (0)...(20)

<223> Plk

158
Gly Leu Glu Leu Ser Arg Arg Arg Ser Leu Leu Glu Leu His Lys Arg
1 5 10 15
Arg Lys Ala Leu
20

159
<210> 159
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> Ret

<400> 159

Gly Val Glu Tyr Ala Lys Tyr Gly Ser Leu Arg Gly Phe Leu Arg Glu
1 5 10 15
Ser Arg Lys Val Gly Pro
20

<210> 160
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)
<223> benzyl ester at position 9

<221> AMIDATION
<222> (0)...(15)

<223> Ret

<400> 160

Gly Ser Leu Arg Gly Phe Leu Arg Glu Ser Arg Lys Val Gly Pro
1 5 10 15

<210> 161
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(21)

<223> Ron

<400> 161

Gly Leu Pro Tyr Met Cys His Gly Asp Leu Leu Gln Phe Ile Arg Ser
1 5 10 15
Pro Gln Arg Asn Pro
20

<210> 162
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(20)

<223> SNK

<400> 162

Gly	Leu	Glu	Tyr	Ser	Ser	Arg	Arg	Ser	Met	Ala	His	Ile	Leu	Lys	Ala
1				5					10					15	
Arg	Lys	Val	Leu												20

<210> 163

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(20)

<223> Syk

<400> 163

Gly	Met	Glu	Met	Ala	Glu	Leu	Gly	Pro	Leu	Asn	Lys	Tyr	Leu	Gln	Gln
1				5					10					15	
Asn	Arg	His	Val												20

<210> 164

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(19)

<223> TGF β RII

<400> 164

Gly	Thr	Ala	Phe	His	Ala	Lys	Gly	Asn	Leu	Gln	Glu	Tyr	Leu	Thr	Arg
1				5					10					15	
His	Val	Ile													

<210> 165

<211> 25

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION
<222> (0)...(25)

<223> TrkB

<400> 165

Gly Phe Glu Tyr Met Lys His Gly Asp Leu Asn Lys Phe Leu Arg Ala
1 5 10 15
His Gly Pro Asp Ala Val Leu Met Ala
20 25

<210> 166

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION

<222> (0)...(13)

<223> TrkB

<400> 166

Gly Leu Arg Ala His Gly Pro Asp Ala Val Leu Met Ala
1 5 10

<210> 167

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION

<222> (0)...(11)

<223> TrkB

<400> 167

Gly Leu Arg Ala His Gly Pro Asp Ala Val Leu
1 5 10

<210> 168

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> TrkB

<400> 168
Gly Leu Asn Phe Lys Leu Arg Ala His Gly Pro Asp Ala
1 5 10

<210> 169
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> TrkB

<400> 169
Gly Phe Lys Leu Arg Ala His Gly Pro Asp Ala Val Leu
1 5 10

<210> 170
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(21)

<223> Zap70

<400> 170
Gly Met Glu Met Ala Gly Gly Gly Pro Leu His Lys Phe Leu Val Gly
1 5 10 15
Lys Arg Glu Glu Ile
20

<210> 171
<211> 21
<212> PRT
<213> Unknown

<220>
<223> IRK

<400> 171

Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu
 1 5 10 15
 Ala Glu Asn Asn Pro
 20

.<210> 172

<211> 8

<212> PRT

<212> FKI

<220>

<223> endothelial growth factor receptor

<400> 172

Lys Phe Asp Val Ile Asn Leu Ala
1 5